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CHEMISTRY

Rapid approach to complex boronic acids

Constantinos G. Neochoritis^{1*†}, Shabnam Shaabani^{1*}, Maryam Ahmadianmoghaddam¹, Tryfon Zarganes-Tzitzikas¹, Li Gao¹, Michaela Novotná¹, Tatiana Mitriková¹, Atilio Reyes Romero¹, Marina Ika Irianti¹, Ruixue Xu¹, Joe Olechno², Richard Ellison², Victoria Helan³, Michael Kossenjans³, Matthew R. Groves¹, Alexander Dömling^{1‡}

The compatibility of free boronic acid building blocks in multicomponent reactions to readily create large libraries of diverse and complex small molecules was investigated. Traditionally, boronic acid synthesis is sequential, synthetically demanding, and time-consuming, which leads to high target synthesis times and low coverage of the boronic acid chemical space. We have performed the synthesis of large libraries of boronic acid derivatives based on multiple chemistries and building blocks using acoustic dispensing technology. The synthesis was performed on a nanomole scale with high synthesis success rates. The discovery of a protease inhibitor underscores the usefulness of the approach. Our acoustic dispensing-enabled chemistry paves the way to highly accelerated synthesis and miniaturized reaction scouting, allowing access to unprecedented boronic acid libraries.

INTRODUCTION

Boron is a unique element of great versatility and individuality, although it seems that nature and evolution have generally bypassed it (with the exception of a few natural products, e.g., boromycin) (1). Boron plays an exquisite role in synthetic chemistry, with boronic acids and their esters of paramount importance to all facets of chemical science. Since the introduction of the Pd-catalyzed C–C Suzuki–Miyaura couplings (2) that brought boronate esters into vogue, the boronic acid moiety has become a very important functional group (3). Other highly useful transformations based on boronic acids include the Petasis reaction (4), C–N and C–O coupling (Chan–Lam coupling) (5, 6), Liebeskind–Srogl coupling (7), regioselective deuteration, or sulfonamide formation (8). Boronic acids as mild electrophiles are also investigated as reversible covalent inhibitors (9, 10), and thousands of different building blocks are now commercially available. As a result, boronic acids are increasingly being seen in approved drugs, e.g., vaborbactam or bortezomib (Fig. 1, A and B) (11, 12).

However, these boron building blocks comprise almost exclusively low-molecular weight compounds, as the late-stage functionalization of high-molecular weight boronic acids is synthetically demanding due to their tedious introduction, modest functional group compatibility, regioselectivity issues, and difficulty to parallelize (13, 14). Because of the exquisite differential properties of boronic acids, an easy access to high-molecular weight elaborated compounds is highly desirable. Isocyanide-based multicomponent reactions (IMCRs) are well established for functional group compatibility that accounts for the immense scaffold diversity that can be generated on the basis of some handful primary IMCRs (15, 16). Furthermore, IMCRs are useful to access a drug-like chemical space and many marketed or experimental drugs (17, 18). Thus, we hypothesized that unprotected boronic acids are compatible with the reaction conditions of IMCR and can be introduced into complex high-molecular weight compounds of use (19). The use of unprotected boronic acids directly

could enable a faster access with limited protecting steps to a large number of boron-based derivatives. In addition, the screening of these compounds (e.g., as covalent inhibitors) could be performed directly

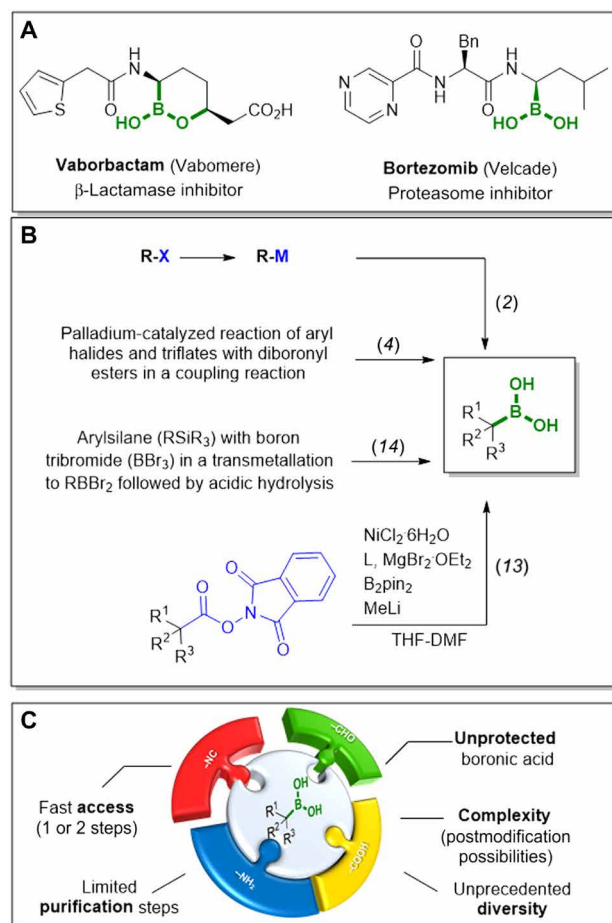


Fig. 1. Importance of boronic acids, commonly used synthetic methods for the $-\text{B}(\text{OH})_2$ introduction, and our proposed building block-centered approach. (A) Marketed drugs containing free $-\text{B}(\text{OH})_2$ moieties. (B) Common methods for late-stage introduction of the $-\text{B}(\text{OH})_2$ moiety. THF-DMF, tetrahydrofuran-dimethylformamide. (C) Building block approach to prepare complex $-\text{B}(\text{OH})_2$ moiety containing molecules in large numbers.

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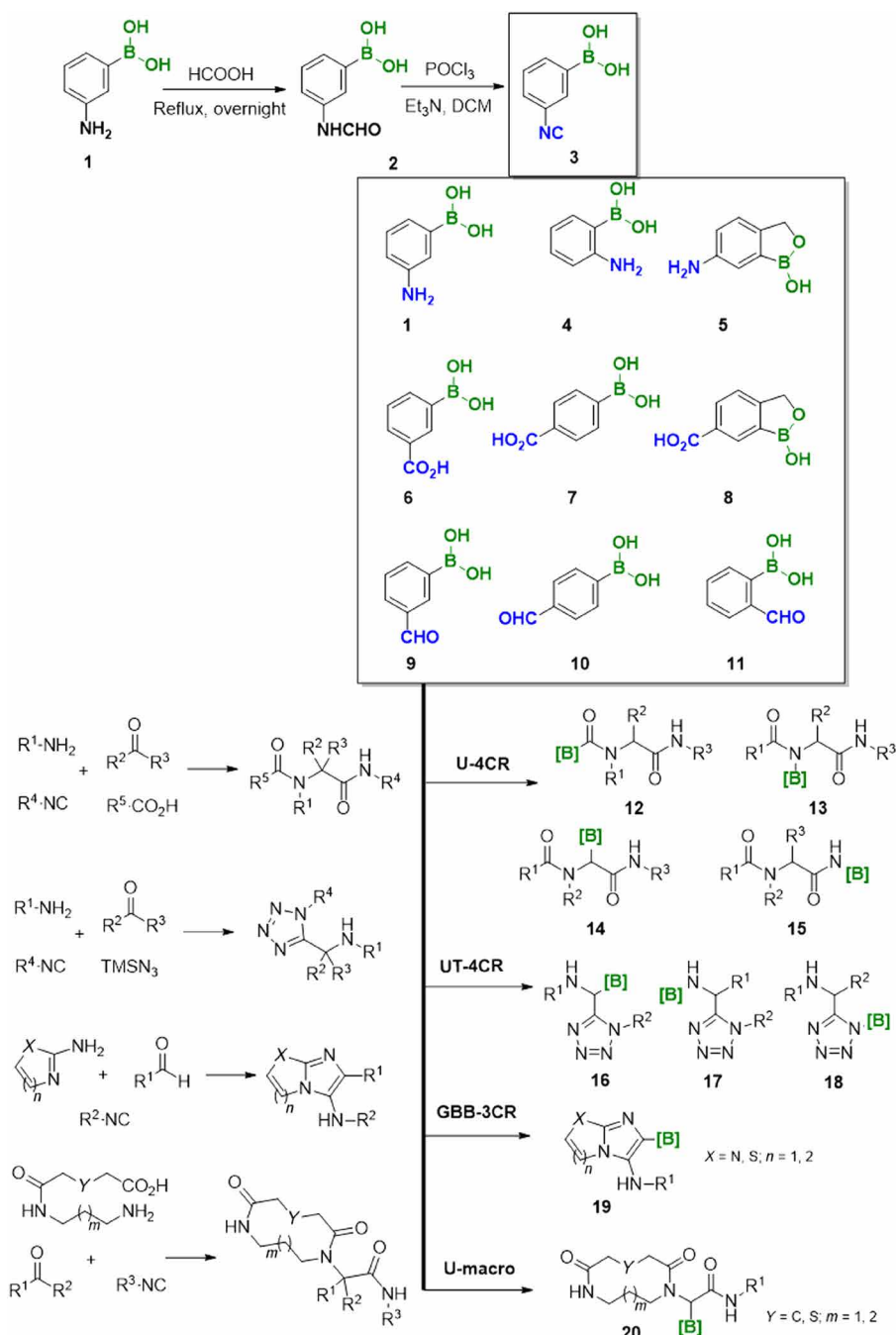


Fig. 2. Boronic acid building blocks used in this study, first synthesis of boronic acid isocyanide, and evaluated reactions. [B], phenyl boronic acid moiety.

without further deprotection (Fig. 1C). To test this hypothesis, we used an acoustic droplet ejection (ADE)-enabled synthesis platform. In ADE, acoustic waves are applied to eject nanoliter droplets from a source plate with building block stock solutions to a destination plate in which the reaction occurs. While ADE is an established dispensing technology in many other scientific areas (e.g., crystallography), it is uncommon in organic synthesis (20). The ADE platform is based on microliter volume chemistry, uses minimal resources, is highly automatable, and is useful to screen many building block combinations in a shorter time frame than other current technologies (21).

RESULTS AND DISCUSSIONS

The mechanism-based functional groups required in IMCRs are carboxylic acids, amines, oxo components, and isocyanides. We synthesized and purchased a number of the first three building block categories. In addition, we also synthesized an unknown isocyanide boronic acid in one example (Fig. 2). We planned to perform four IMCRs and investigate the reaction success rate depending on the reactions, the components, and the substitution pattern (e.g., *o*-, *m*-, and *p*-) in combination with multiple complementary building blocks chosen in a random fashion.

Optimizing the compatibility of boronic acids for MCR is an interesting synthetic challenge, as the C—B bond is well known to react

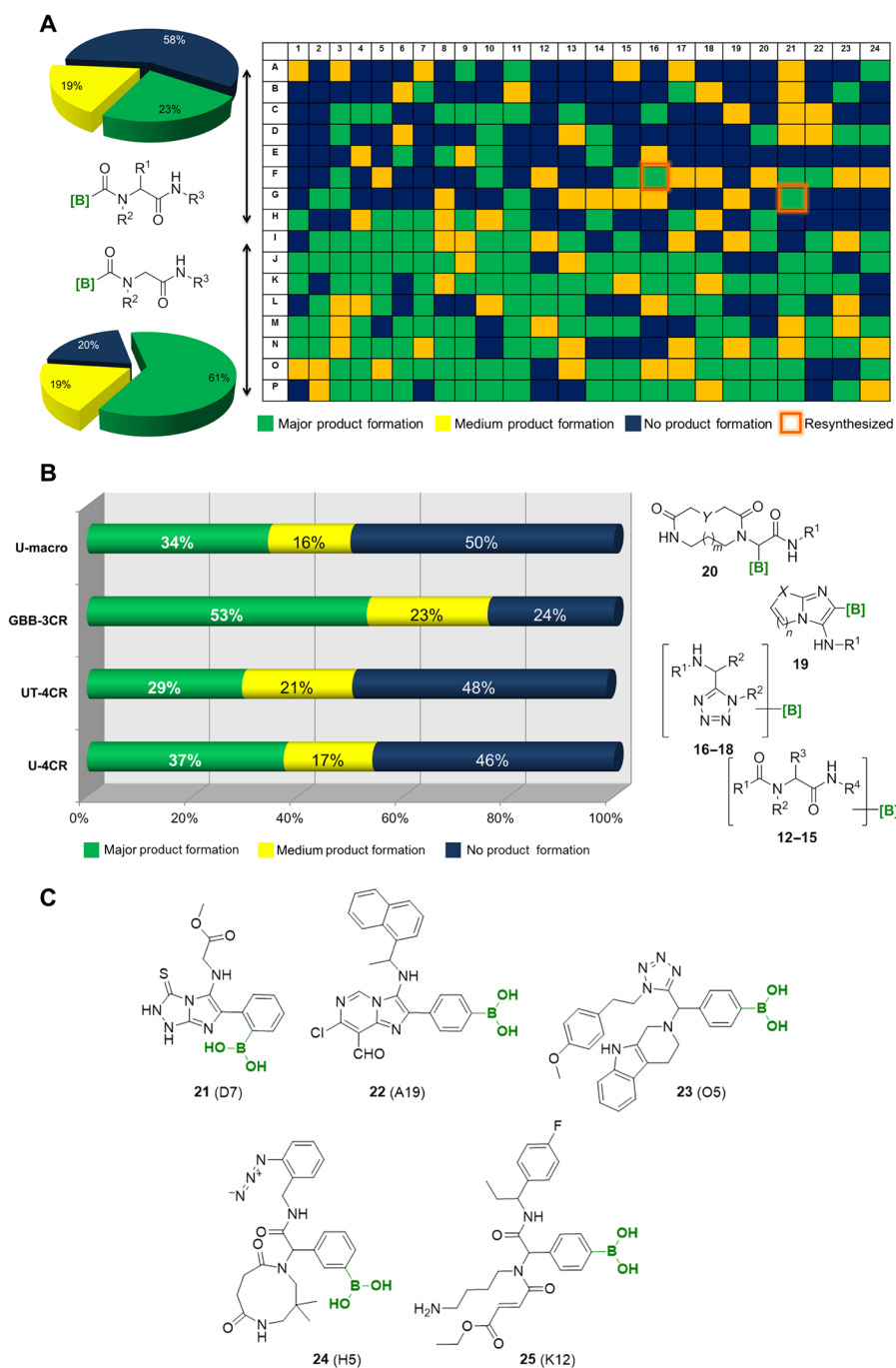


Fig. 3. HT synthesis of boronic acids using the building block approach. (A) Exemplary analytical 384-well plate of the U-4CR scaffold **12** (green, major product formation; yellow, product present; blue, product not present). (B) Statistical analysis of the quality of reactions of the different scaffolds. (C) Structures of some unusual reaction products from different IMCRs.

with common MCR starting materials and intermediates under mild conditions, e.g., primary, secondary amines, and carbonyl compounds (Petasis reaction and others) (22). Moreover, the electrophilic boron could unproductively complex to nucleophilic key functional groups of MCRs and thereby interrupt the reaction progress (23). Here, we used different building blocks (amines, aldehydes, carboxylic acids, and isocyanides) with free boronic acids in different positions to

investigate their compatibility with a number of IMCRs (Fig. 2). The stability of the boronic acid moiety in the presence of the isocyanide in one molecule in the absence of such molecules is unknown. To this end, we also synthesized the first free boronic acid-containing isocyanide. We have extensively investigated the compatibility of multiple free boronic acid-containing building blocks in multiple IMCRs, including the classical Ugi four-component reaction (U-4CR) (24),

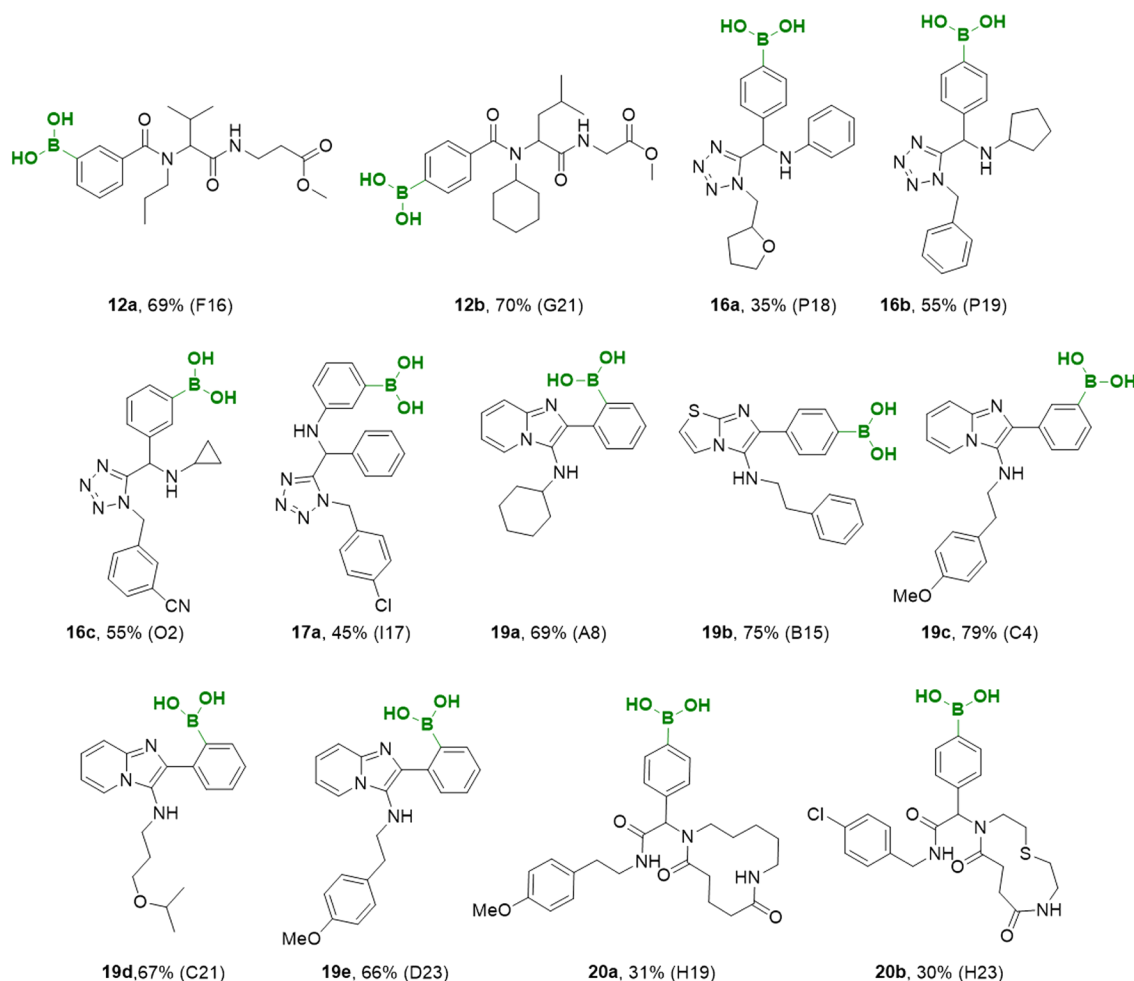


Fig. 4. Resynthesized complex boronic acid derivatives based on different scaffolds on a millimole scale and corresponding yields.

the Ugi tetrazole (UT-4CR) (25), the Gröbcke-Blackburn-Bienaymé (GBB-3CR) (26–28), and the Ugi-based macrocycles (scaffolds **12** to **20**; Fig. 3) (29). In addition, we studied the suitability of the corresponding boronic acid libraries in a secondary Suzuki cross-coupling reaction. We performed the project under extreme resource- and time-saving conditions using the ADE-enabled chemistry platform in a 384-well format.

We accomplished the synthesis of *m*-isocyanophenyl boronic acid by a classical formylation/dehydration procedure (Fig. 2). Because of the presumed instability, we immediately used the isocyanide after preparation. Using the boronic acid building blocks of Fig. 2, we investigated different IMCRs on a nanomole scale using ADE technology. The analytics of the four 384-well format plates were performed using mass spectrometry (MS) as described previously (30), which allowed us to classify the reactions into three groups: major (green color), mediocre (yellow color), or no product formation (blue color). The outcome of the high-throughput (HT) analytics for the different reactions is shown in Fig. 3, and a detailed analysis of the different building blocks is given in the Supplementary Materials. The rapid collection of information facilitated the ability to predict outcomes from other possible combinations of reagents. For example, it was found that the *ortho* substituted building blocks **4** and **11** reacted less efficiently than the corresponding *meta* or *para* substi-

tuted in all MCRs (Fig. 2). This can be rationalized by the neighbor group effect of boronic acid that might hamper formation or reduce reactivity of the key Schiff base. It was also found that boronic acid monoesters **5** and **8** were less reactive than their boronic acid counterparts, probably due to the introduction of ring strain around the boron center, leading to slightly different electronic properties (31). In addition, it was demonstrated that the U-4CR of the three carboxyphenyl boronic acids **6** to **8** (heat map shown in Fig. 3A) was greatly enhanced when *p*-formaldehyde was used (>60% of the reactions worked; see the Supplementary Materials). Last, it is noteworthy that formylphenyl boronic acids behave well in the GBB-3CR, since more than 50% of the reactions that were performed were successful (see the Supplementary Materials). In general, the use of building blocks without the free —B(OH)_2 moieties was less successful than those with boronic acids. This could point to a potential catalytic activity of boronic acids in the GBB-3CR as a Brønsted acid as there are many cases of GBB catalysis by Brønsted acids (26–28). Detailed analysis of the rich data of the complementary building blocks can help to uncover subtle reactivity details.

In our approach, novel substrates could be generated. In the GBB-3CR, compound **21** reacted repeatedly well despite the existence of a hitherto unreported triazolidine-5-thione moiety in this context. Another interesting finding is the good reactivity of building

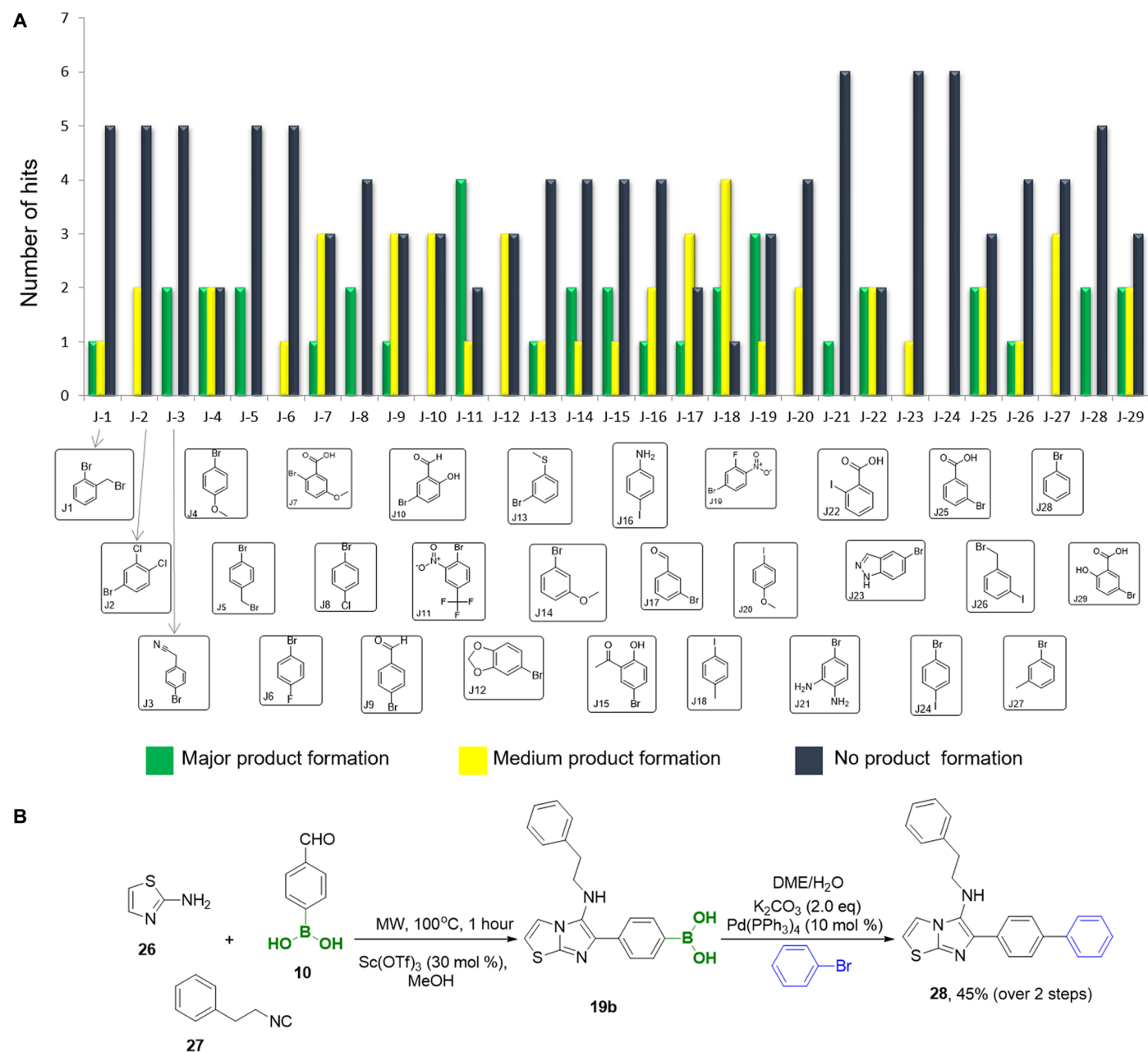


Fig. 5. HT Suzuki reaction of boronic acids using the building block approach. (A) Statistical analysis of the aryl halides that were used in the HT screening (green, major peak in MS; yellow, product present; blue, product not present). (B) A one-pot resynthesized compound **28** on a millimole scale and isolated yield. DME, dimethoxyethane.

block **22** in the GBB-3CR, in which the formyl group did not react, as the additional formyl group could theoretically undergo alternative reaction pathways such as condensation and addition reactions. Another pleasant finding is the good reactivity of tetrahydro- β -carboline **23** in the UT-4CR, which is a pharmacophore in multiple natural products and drugs (e.g., harman and tadalafil). Complex medium-sized and macrocycles gave, unexpectedly, very good results (e.g., medium-sized cycle **24**). Last, we observed functional group tolerance and

selectivity. In the case of U-4CR **25**, we used a diamine, which reacted only once, leaving a primary amine behind. The HT synthesis approach displayed here is a treasure trove to uncover interesting unknown reactivities that deserve further investigation and detailed analysis in a narrower compound series.

The scalability from the nanomole to the millimole scale is often problematic. Therefore, we resynthesized multiple examples of each compound series (compounds **12** to **20**) on a millimole scale (including

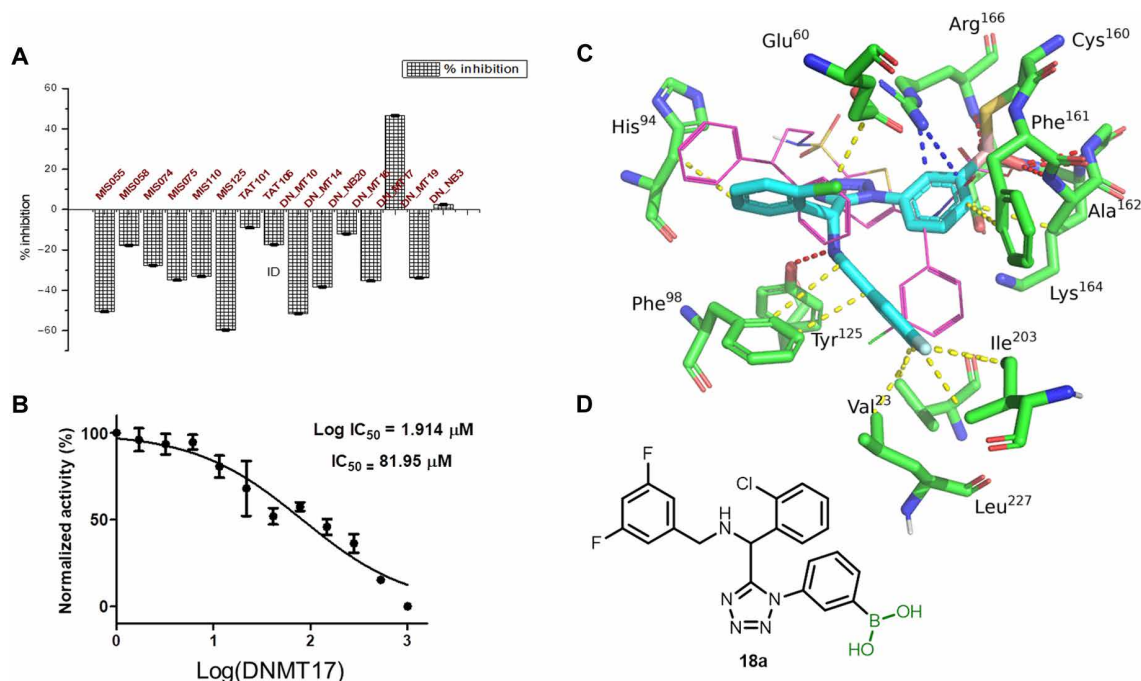


Fig. 6. Covalent inhibition of tuberculosis target MptpB. (A) Screening of the boronic acid library by a colorimetric enzyme assay. (B) Median inhibitory concentration (IC_{50}) of compound **18a**. (C and D) Modeling of compound **18a** into MptpB [Protein Data Bank (PDB) ID: 2OZ5], where it forms a covalent adduct with active-site cysteine. Van der Waals interactions, hydrogen bonding, and cation- π interactions are indicated by yellow, red, and blue dotted lines, respectively.

full characterization) to verify our ADE results (Fig. 4) and to identify any potential bottlenecks into transferring synthesis from ADE technology to classical approaches.

Boronic acids are exceedingly useful functional groups and the starting materials for many reactions. To underscore the usefulness of our building block approach, we investigated multiple boronic acid building blocks in a subsequent Suzuki C—C coupling. We performed reaction scouting in 384-well polypropylene plates using ADE; after the transfer of starting materials, we incubated the plates at 50°C overnight. Again, we resynthesized an example on a millimole scale in a one-pot fashion (compound **28**; Fig. 5).

To further underscore the usefulness of our fast, convergent, and highly diverse access of boronic acid libraries, we screened for inhibition of the biological target MptpB, a virulence factor from *Mycobacterium tuberculosis* (32). MptpB belongs to the notoriously undruggable target class of phosphatases that, despite their overarching relevance in medicine, suffer from having no approved drug (33). This is generally attributed to the highly positively charged active site of phosphatases requiring negatively charged inhibitors that cannot overcome membrane penetration issues (34). Looking for a potential covalent interaction between the active-site nucleophiles Cys¹⁶⁰, Thr²²³, and Ser⁵⁷ of MptpB and an electrophilic boronic acid, we screened the library in a colorimetric enzyme assay (see the Supplementary Materials). In this assay, we found several hits, the most potent one **18a** (Fig. 6). The exact binding mode of **18a** is unclear due to the large number of reactive Ser, Cys, and Thr on the surface and in the active site of MptpB (the Supplementary Materials). Modeling studies of **18a** in MptpB with Cys¹⁶⁰, Ser⁵⁷, and Thr²²³ were performed and suggest a covalent adduct to a tetrahedral boron (Fig. 6C and see the Supplementary Materials).

Classical access to boronic acids by late-stage functionalization of complex molecules suffers from a lack in functional group com-

patibility and regioselectivity and often requires harsh conditions that are incompatible with molecule stability (35–38). Here, we introduced the concept of boronic acid building blocks combined with the diversity of MCRs as a valid approach for the synthesis of large and unprecedented libraries of boronic acids. Our studies go much beyond a singleton report on the use of a few free boronic acids in the Ugi reaction as we also investigated the GBB-3CR, UT-4CR, and several different IMCR variations more in an unprecedented breadth of building block combinations (35, 36). In other reports, isocyanide-bearing boronic acids are only known in their protected ester form that would need another, often harsh, deprotecting step to yield boronic acids suitable for screening (39, 40). Here, we found that IMCR generally runs under such mild conditions that free boronic acids are widely tolerated. We systematically investigated 10 different boronic acid building blocks with complementary functional groups (primary amine, aldehyde, and isocyanide) and combined them with 353 different reactants in four IMCRs. More than 1300 different combinations were investigated in a nanomole miniaturized and automated fashion using ADE technology. HT analytics using MS revealed that the different reactions worked better than satisfactorily in 714 cases (458 giving the main product and 256 cases a satisfactory yield). Many subtle reactivities were uncovered, which in a classical millimole scale reaction, evaluation approach could never been elucidated in a reasonable time frame. Upscaling of a substantial number of diverse products revealed the synthetic usefulness of the approach. Last, we probed our library to uncover previously unknown boronic acid-based covalent inhibitors for a notoriously undruggable phosphatase target, identifying a micromolar inhibitor. We believe our described building block approach will widen the accessibility of the boronic acid chemical space markedly for applications in synthesis, chemical biology, and drug discovery. This is also true in light of the recently found catalytic enantioselective Ugi reaction (41).

MATERIALS AND METHODS

All the reagents and solvents were purchased from Sigma-Aldrich, AK Scientific, Fluorochem, abcr GmbH, and Acros and were used without further purification. All isocyanides were prepared in-house (see the Supplementary Materials). All microwave irradiation reactions were carried out in a Biotage Initiator microwave synthesizer. Thin-layer chromatography was performed on Millipore precoated silica gel plates (thickness, 0.20 mm; particle size, 25 μ m). Nuclear magnetic resonance spectra were recorded on Bruker Avance 500 spectrometers [^1H NMR (nuclear magnetic resonance; 500 MHz), ^{13}C NMR (126 MHz)]. Chemical shifts for ^1H NMR were reported as δ values, and coupling constants were in hertz. The following abbreviations were used for spin multiplicity: s, singlet; br s, broad singlet; d, doublet; t, triplet; q, quartet; quin, quintet; dd, double of doublets; ddd, double doublet of doublets; and m, multiplet. Chemical shifts for ^{13}C NMR were reported in parts per million relative to the solvent peak. Flash chromatography was performed on a Reveleris X2 flash chromatography system, using Grace Reveleris Silica flash cartridges (12 g). Mass spectra were measured on a Waters Investigator Supercritical Fluid Chromatograph with a 3100 MS detector (ESI) using a solvent system of methanol and CO_2 on a Viridis silica gel column (4.6 \times 250 mm, 5- μ m particle size) or Viridis 2-ethyl pyridine column (4.6 \times 250 mm, 5- μ m particle size). High-resolution mass spectra were recorded using an LTQ Orbitrap XL (Thermo Fisher Scientific) at a resolution of 60,000 at m/z 400. The Echo 555 liquid handler (Labcyte) was used to transfer nanoliter droplets of starting materials from the 384-well source plate to the 384-well destination plate.

SUPPLEMENTARY MATERIALS

Supplementary material for this article is available at <http://advances.sciencemag.org/cgi/content/full/5/7/eaaw4607/DC1>

Supplementary Materials and Methods

Fig. S1. Isocyanide syntheses.

Fig. S2. Reactions in destination plate I.

Fig. S3. Reactions in destination plate II.

Fig. S4. Reactions in destination plate III.

Fig. S5. Reactions in destination plate IV.

Fig. S6. Labcyte Echo plate reformat software.

Fig. S7. Heat plots with product structures, green for major product formation, yellow for medium product formation, and blue for no product formation.

Fig. S8. Stabilization effect of **18a** as proof of interaction with MptpB as assessed by DSF.

Fig. S9. Binding curve of **18a** to the fluorescently labeled MptpB sample as assessed by MST.

Fig. S10. Three-dimensional structure of the target phosphatase.

Fig. S11. Proposed docking model for **18a** covalently bound to Cys¹⁶⁰ (PDB ID: 2OZ5).

Fig. S12. Proposed docking model for **18a** covalently bound to Ser⁵⁷ (PDB ID: 2OZ5).

Fig. S13. Proposed docking model for **18a** covalently bound to Thr²²³ (PDB ID: 2OZ5).

Fig. S14. ADE technology.

Table S1. Summary table of the docking scores for Covdock and Scorpion.

Scheme S1. Quality control results for destination plate I.

Scheme S2. Performance of formylphenyl boronic acids in destination plate I.

Scheme S3. Performance of isocyanides in GBB-3CR reaction in destination plate I.

Scheme S4. Performance of isocyanides in Ugi-based macrocycles in destination plate I.

Scheme S5. Performance of isocyanides in U-4CR in destination plate I.

Scheme S6. Performance of isocyanides in UT-4CR in destination plate I.

Scheme S7. Performance of carboxylic acids in U-4CR in destination plate I.

Scheme S8. Performance of amines in U-4CR in destination plate I.

Scheme S9. Performance of amines in UT-4CR in destination plate I.

Scheme S10. Performance of amidines in GBB-3CR reaction in destination plate I.

Scheme S11. Performance of α,ω -amino carboxylic acids in Ugi-based macrocycles in destination plate I.

Scheme S12. Quality control results for destination plate II.

Scheme S13. Performance of aminophenyl boronic acids in destination plate II.

Scheme S14. Performance of isocyanides in U-4CR in destination plate II.

Scheme S15. Performance of isocyanides in UT-4CR in destination plate II.

Scheme S16. Performance of oxo components in U-4CR in destination plate II.

Scheme S17. Performance of oxo components in UT-4CR in destination plate II.

Scheme S18. Performance of carboxylic acids in U-4CR in destination plate II.

Scheme S19. Quality control results for destination plate III.

Scheme S20. Performance of carboxyphenyl boronic acids in destination plate III.

Scheme S21. Performance of isocyanides in U-4CR in destination plate III.

Scheme S22. Performance of isocyanides in U-4CR with CH_2O in destination plate III.

Scheme S23. Performance of oxo component in U-4CR in destination plate III.

Scheme S24. Performance of amines in U-4CR in destination plate III.

Scheme S25. Performance of amines in U-4CR with CH_2O in destination plate III.

Scheme S26. Quality control results for destination plate IV.

Scheme S27. Performance of MCR boronic acid building blocks in destination plate IV.

Scheme S28. Performance of aryl halides in destination plate IV.

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